

AMENDMENT TO THE SPECIFICATION

On pages 16-20, please amend Table 2 as shown on pages 3-7 of this Supplemental Response.

On pages 159-162, please amend Table 11 as shown on pages 8-11 of this Supplemental response.

TABLE 2

CRY3BB* PROTEINS EXHIBITING IMPROVED ACTIVITY AGAINST SCRWF LARVAE

Cry3Bb*	cry3Bb*	cry3Bb* Nucleotide Sequence	Cry3Bb* Amino Acid Changes	Structural Site of Changes	Fold Increase Over WT Activity	Design Method Used
Protein	Plasmid	Changes				
Designation	Designation					
Cry3Bb.60	—	—	Δ1-159	Δα1-α3	3.6×	1, 6, 8
Cry3Bb.11221	pEG1707	A460T, C461T, A462T, C464A, T465C, T466C, T467A, A468T, A469T, G470C, T472C, T473G, G474T, A477T, A478T, G479C	T154F, P155H, L156H, L158R	lα3,4	6.4×	1, 8
Cry3Bb.11222	pEG1708	T687C, T688C, A689T, C691A, A692G	Y230L, H231S	α6	4.0×	3, 7
Cry3Bb.11223	pEG1709	T667C, T687C, T688A, A689G, C691A, A692G	S223P, Y230S	α6	2.8×	3
Cry3Bb.11224	pEG1710	T687C, A692G	H231R	α6	5.0×	7, 8
Cry3Bb.11225	pEG1711	T687C, C691A	H231N, T241S	α6	3.6×	7
Cry3Bb.11226	pEG1712	T687C, C691A, A692C, T693C	H231T	α6	3.0×	7, 8
Cry3Bb.11227	pEG1713	C868A, G869A, G870T	R290N	lα7,β1	1.9×	2, 3, 4, 6
Cry3Bb.11228	pEG1714	C932T, A938C, T942G, G949A, T954C	S311L, N313T, E317K	lβ1,α8	4.1×	2, 4

Cry3Bb*	<i>cry3Bb*</i>	<i>cry3Bb*</i> Nucleotide Sequence	Cry3Bb* Amino Acid Changes	Structural Site of Changes	Fold Increase Over WT Activity	Design Method Used
Protein	Plasmid	Changes				
Designation	Designation					
Cry3Bb.11229	pEG1715	T931A, A933C, T942A, T945A, G949A, A953G, T954C	S311T, E317K, I β 1, α 8	2.5 \times	2, 4	
Cry3Bb.11230	pEG1716	T931G, A933C, C934G, T945G, C946T, A947G, G951A, T954C	S311A, L312V, Q316W	I β 1, α 8	4.7 \times	2, 4, 8
Cry3Bb.11231	pEG1717	T687C, A692G, C932T, A938C, T942G, G949A, T954C	H231R, S311L, α 6; I β 1, α 8	I β 1, α 8	7.9 \times	2, 4, 7, 8, 10
Cry3Bb.11232	pEG1718	T931A, A933G, T935C, T936A, A938C, T939C, T942C, T945A, G951T, T954C	N313T, E317K, S311T, L312P, N313T, E317N	I β 1, α 8	5.1 \times	4
Cry3Bb.11233	pEG1719	T931G, A933C, T936G, T942C, C943T, T945A, C946G, G948C, T954C	S311A, Q316D	I β 1, α 8	2.2 \times	2, 4
Cry3Bb.11234	pEG1720	T861C, T866C, C868A, T871C, T872G, A875T, T877A, C878G, A882G	I289T, L291R, Y292F, S293R	I α 7, β 1	4.1 \times	4
Cry3Bb.11235	pEG1721	T687C, A692G, C932T	H231R, S311L	α 6; I β 1, α 8	3.2 \times	2, 4, 7, 8, 10

Cry3Bb*	<i>cry3Bb*</i>	<i>cry3Bb*</i> Nucleotide Sequence	Cry3Bb* Amino Acid Changes	Structural Site of Changes	Fold Increase Over WT Activity	Design Method Used
Protein	Plasmid	Changes				
Designation	Designation					
Cry3Bb.11236	pEG1722	T931A, C932T, A933C, T936C, T942G, T945A, T954C	S311I	I β 1, α 8	3.1 \times	2, 4
Cry3Bb.11237	pEG1723	T931A, C932T, A933C, T936C, A937G, A938T, C941A, T942C, T945A, C946A, A947T, A950T, T954C	S311I, N313H	I β 1, α 8	5.4 \times	2, 4
Cry3Bb.11238	pEG1724	A933C, T936C, A937G, A938T, C941A, T942C, T945A, C946A, A947T, A950T, T954C	N313V, T314N, Q316M, E317V	I β 1, α 8	2.6 \times	2, 4
Cry3Bb.11239	pEG1725	A933T, A938G, T939G, T942A, T944C, T945A, A947T, G948T, A950C, T954C	N313R, L315P, Q316L, E317A	I β 1, α 8	2.8 \times	2, 4
Cry3Bb.11241	pEG1726	A860T, T861C, G862A, C868T, G869T, T871C, A873T, T877A, C878G, A879T	Y287F, D288N, R290L	I α 7, β 1	2.6 \times	2, 3, 4, 6
Cry3Bb.11242	pEG1727	C868G, G869T	R290V	I α 7, β 1	2.5 \times	2, 3, 4, 6, 8

Cry3Bb*	<i>cry3Bb*</i>	<i>cry3Bb*</i> Nucleotide Sequence	Cry3Bb* Amino Structural Site	Fold Increase Over WT Activity	Design
Protein	Plasmid	Changes	of Changes	WT Activity	Method Used
Designation	Designation				
Cry3Bb.11032	pEG1041	A494G	D165G	3.1x	2, 4, 8
Cry3Bb.11035	pEG1046	G479A, A481C, A482C, A484C, G485A, A486C, A494G	S160N, K161P, <u>P</u> R162H, D165G	2.7x	8
Cry3Bb.11036	pEG1047	A865G, T877C	I289V, S293P	4.3x	4
Cry3Bb.11046	pEG1052	G479A, A481C, A482C, A484C, G485A, A486C, A494G, A865G, T877C	S160N, K161P, <u>P</u> R162H, D165G, I289V, S293P	2.6x	2, 4, 8, 10
Cry3Bb.11048	pEG1054	T309A, Δ 310, Δ 311, Δ 312	D103E, Δ A104	4.3x	8
Cry3Bb.11051	pEG1057	A565G, A566G	K189G	3.0x	2, 3, 4
Cry3Bb.11057	pEG1062	T309A, Δ 310, Δ 311, Δ 312, G479A, A481C, A482C, A484C, G485A, A486C, A494G	D103E, Δ A104, S160N, K161P, <u>P</u> R162H, D165G	3.4x	2, 4, 8, 10
Cry3Bb.11058	pEG1063	T309A, Δ 310, Δ 311, Δ 312, A460T, C461T, A462T, C464A, T465C, T466C, T467A, A468T, A469T, G470C, T472C, T473G, G474T, A477T, A478T, G479C	D103E, Δ A104, Δ 2a,2b; Δ 3,4	3.5x	1, 8, 10

Designation	Protein	Plasmid	<i>cry3Bb*</i>	<i>cry3Bb*</i> Nucleotide Sequence	Cry3Bb* Amino Acid Changes	Structural Site of Changes	Fold Increase Over WT Activity	Design Method Used
Cry3Bb.11081	pEG1084		A494G, T931A, A933C, T942A, T945A, G949A, T954C	D165G, S311T, E317K	$\alpha 4; \beta 1, \alpha 8$	$6.1 \times$	$2, 4, 8, 10$	
Cry3Bb.11082	pEG1085		A494G, A865G, T877C, T914C, T931G, A933C, C934G, T945G, C946T, A947G, G951A, T954C, A1043G, T1094C	D165G, I289V, S293P, F305S, S311A, L312V, Q316W, Q348R, V365A	$\alpha 4; \beta 1, \beta 1; \beta 2; \beta 3b$	$4.9 \times$	$2, 4, 5, 8, 9, 10$	
Cry3Bb.11083	pEG1086		A865G, T877C, A1043G	I289V, S293P, Q348R	$\beta 1, \beta 1; \beta 2$	$7.4 \times$	$4, 5, 9, 10$	
Cry3Bb.11084	pEG1087		A494G, C932T	D165G, S311L	$\alpha 4; \beta 1, \alpha 8$	$7.2 \times$	$2, 4, 8, 10$	
Cry3Bb.11095	pEG1095		A1043G	Q348R	$\beta 2$	$4.6 \times$	$5, 9$	
Cry3Bb.11098	pEG1098		A494G, T687C, A692G, C932T, A938C, T942G, G949A, T954C	D165G, H231R, S311L, N313T, E317K	$\alpha 4; \alpha 6, 1\beta 1, \alpha 8$	$7.9 \times$	$2, 4, 7, 8$	

TABLE 11

DNA SEQUENCE CHANGES OF CRY3BB* GENES AND RESULTING AMINO ACID SUBSTITUTIONS OF THE CRY3BB* PROTEINS

Plasmid	<i>cry3Bb*</i> DNA Sequence	Cry3Bb* Amino Acid Sequence	Structural Site of Alteration
pEG1707	A460T, C461T, A462T, C464A, T465C, T466C, T467A, A468T, A469T, G470C, T472C, T473G, G474T, A477T, A478T, G479C	T154F, P155H, L156H, L158R	$\alpha 3,4$
pEG1708	T687C, T688C, A689T, C691A, A692G	Y230L, H231S	$\alpha 6$
pEG1709	T667C, T687C, T688A, A689G, C691A, A692G	S223P, Y230S	$\alpha 6$
pEG1710	T687C, A692G	H231R	$\alpha 6$
pEG1711	T687C, C691A	H231N, T241S	$\alpha 6$
pEG1712	T687C, C691A, A692C, T693C	H231T	$\alpha 6$
pEG1713	C868A, G869A, G870T	R290N	$\alpha 7, \beta 1$
pEG1714	C932T, A938C, T942G, G949A, T954C	S311L, N313T, E317K	$\beta 1, \alpha 8$
pEG1715	T931A, A933C, T942A, T945A, G949A, A953G, T954C	S311T, E317K, Y318C	$\beta 1, \alpha 8$
pEG1716	T931G, A933C, C934G, T945G, C946T, A947G, G951A, T954C	S311A, L312V, Q316W	$\beta 1, \alpha 8$
pEG1717	T687C, A692G, C932T, A938C, T942G, G949A, T954C	H231R, S311L, N313T, E317K	$\alpha 6, \beta 1, \alpha 8$

Plasmid	<i>cry3Bb*</i> DNA Sequence	Cry3Bb* Amino Acid Sequence	Structural Site of Alteration
pEG1718	T931A, A933G, T935C, T936A, A938C, T939C, T942C, T945A, G951T, T954C	S311T, L312P, N313T, E317N	β 1, α 8
pEG1719	T931G, A933C, T936G, T942C, C943T, T945A, C946G, G948C, T954C	S311A, Q316D	β 1, α 8
pEG1720	T861C, T866C, C868A, T871C, T872G, A875T, T877A, C878G, A882G	I289T, L291R, Y292F, S293R	α 7, β 1
pEG1721	T687C, A692G, C932T	H231R, S311L	α 6, β 1, α 8
pEG1722	T931A, C932T, A933C, T936C, T942G, T945A, T954C	S311I	β 1, α 8
pEG1723	T931A, C932T, A933C, T936C, A937G, A938T, C941A, T942C, T945A, C946A, A947T, A950T, T954C	S311I, N313H	β 1, α 8
pEG1724	A933C, T936C, A937G, A938T, C941A, T942C, T945A, C946A, A947T, A950T, T954C	N313V, T314N, Q316M, E317V	β 1, α 8
pEG1725	A933T, A938G, T939G, T942A, T944C, T945A, A947T, G948T, A950C, T954C	N313R, L315P, Q316L, E317A	β 1, α 8
pEG1726	A860T, T861C, G862A, C868T, G869T, T871C, A873T, T877A, C878G, A879T	Y287F, D288N, R290L	α 7, β 1
pEG1727	C868G, G869T	R290V	α 7, β 1

Plasmid	<i>cry3Bb</i> * DNA Sequence	Cry3Bb* Amino Acid Sequence	Structural Site of Alteration
pEG1041		A494G	D165G
pEG1046	G479A, A481C, A482C, A484C, G485A, A486C, A494G	S160N, K161P, <u>P</u> R162H, D165G	α 4
pEG1047	A865G, T877C	I289V, S293P	α 7, β 1
pEG1052	G479A, A481C, A482C, A484C, G485A, A486C, A494G, A865G, T877C	S160N, K161P, <u>P</u> R162H, D165G, I289V, S293P	α 4, α 7, β 1
pEG1054	T309A, Δ 310, Δ 311, Δ 312	D103E, Δ A104	α 2a, 2b
pEG1057	A565G, A566G	K189G	α 4, 5
pEG1062	T309A, Δ 310, Δ 311, Δ 312, G479A, A481C, A482C, A484C, G485A, A486C, A494G	D103E, Δ A104, S160N, K161P, <u>P</u> R162H, D165G	α 2a, 2b α 4
pEG1063	T309A, Δ 310, Δ 311, Δ 312, A460T, C461T, A462T, C464A, T465C, T466C, T467A, A468T, A469T, G470C, T472C, T473G, G474T, A477T, A478T, G479C	D103E, Δ A104, T154F, P155H, L156H, L158R	α 2a, 2b α 3, 4
pEG1084	A494G, T931A, A933C, T942A, T945A, G949A, T954C	D165G, S311T, E317K	α 4, β 1, α 8

Plasmid	<i>cry3Bb*</i> DNA Sequence	<i>Cry3Bb*</i> Amino Acid Sequence	Structural Site of Alteration
pEG1085	A494G, A865G, T877C, T914C, T931G, A933C, C934G, T945G, C946T, A947G, G951A, T954C, A1043G, T1094C	D165G, I289V, S293P, F305S, S311A, L312V, Q316W, Q348R, V365A	$\alpha 4, \text{l}\alpha 7, \beta 1, \beta 1, \text{l}\beta 1, \alpha 8$ $\beta 2, \beta 3\text{b}$
pEG1086	A865G, T877C, A1043G	I289V, S293P, Q348R	$\text{l}\alpha 7, \beta 1, \beta 2$
pEG1087	A494G, C932T	D165G, S311L	$\alpha 4, \text{l}\beta 1, \alpha 8$
pEG1095	A1043G	Q348R	$\beta 2$